IN THE DRAWINGS:

Please replace the original Figure 11 with the substitute Figure 11 attached hereto.

Figure 11

SEQ ID NO:33

Nod2 cDNA sequence

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gtagacagat ccaggctcac cagtcctgtg ccactgggct tttggcgttc tgcacaaggc        60

ctacccgcag atgccatgcc tgctccccca gcctaatggg ctttgatggg ggaagagggt       120

ggttcagcct ctcacgatga ggaggaaaga gcaagtgtcc tcctcggaca ttctccgggt       180

tgtgaaatgt gctcgcagga ggcttttcag gcacagagga gccagctggt cgagctgctg       240

gtctcagggt ccctggaagg cttcgagagt gtcctggact ggctgctgtc ctgggaggtc       300

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cacagccatg tggagaacat gctggacctg gcatgggagc ggggtttcgt cagccagtat       600

gaatgtgatg aaatcaggtt gccgatcttc acaccgtccc agagggcaag aaggctgctt       660

gatcttgcca cggtgaaagc gaatggattg gctgccttcc ttctacaaca tgttcaggaa       720

ttaccagtcc cattggccct gcctttggaa gctgccacat gcaagaagta tatggccaag       780

ctgaggacca cggtgtctgc tcagtctcgc ttcctcagta cctatgatgg agcagagacg       840

ctctgcctgg aggacatata cacagagaat gtcctggagg tctgggcaga tgtgggcatg       900

gctggacccc cgcagaagag cccagccacc ctgggcctgg aggagctctt cagcacccct       960

ggccacctca atgacgatgc ggacactgtg ctggtggtgg gtgaggcggg cagtggcaag      1020

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ctctttgtct tcccattcag ctgccggcag ctgcagtgca tggccaaacc actctctgtg      1140

cggactctac tctttgagca ctgctgttgg cctgatgttg gtcaagaaga catcttccag      1200

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ttcaggttca cggatcgtga acgccactgc tccccgaccg accccacctc tgtccagacc      1320
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tcagcccagc agctccaggc agcacaggtc agccctgatg acatttctct tggcttcctg      1860

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cactggggcc tgctggctga gtgccagaca tctgagaagg ccctgctccg gcgccaggcc      2220

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ctccttgcat gcaggcagaa cttcttggca ttgaggctgg ggaataacta catcactgcc      2760

gcgggagccc aagtgctggc cgaggggctc cgaggcaaca cctccttgca gttcctggga      2820

ttctggggca acagagtggg tgacgagggg gcccaggccc tggctgaagc cttgggtgat      2880

caccagagct tgaggtggct cagcctggtg gggaacaaca ttggcagtgt gggtgcccaa      2940

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catctccagg atgaaggtgt atgttctctc gcagaaggac tgaagaaaaa ttcaagtttg      3060

aaaatcctga agttgtccaa taactgcatc acctacctag gggcagaagc cctcctgcag      3120
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Figure 11

SEQ ID NO:33

Nod2 cDNA sequence

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ctacccgcag atgccatgcc tgctcccca gcctaatggg ctttgatggg ggaagagggt          120

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cacagccatg tggagaacat gctggacctg gcatgggagc ggggtttcgt cagccagtat          600

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gatcttgcca cggtgaaagc gaatggattg gctgccttcc ttctacaaca tgttcaggaa          720

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